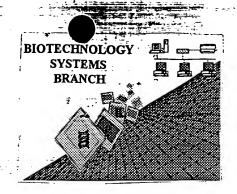
## RAW-SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: \_\_\_\_\_\_\_\_\_

11.48

Date Processed by STIC:

Source:

1/26/2000

JAN 05 7001 TECH CENTER 1600/2960

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

## Raw Sequence Listing Error Summary

## ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/497,997

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1	Wrapped Nucleics	The number/text at the end of each line "wrapped" down to the next line.
		This may occur if your file was retrieved in a word processor after_creating it.
		Please adjust your right margin to .3, as this will prevent "wrapping".
2	Wrapped Aminos	The amino acid number/lext at the end of each line "wrapped" down to the next line.
	_	This may occur if your file was retrieved in a word processor after creating it.
	•	Please adjust your right margin to .3, as this will prevent "wrapping".
3	Incorrect Line Length	The rules require that a line not exceed 72 characters in length. This includes spaces.
. 4	Misaligned Amino Acid	The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs
,	Numbering	between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
5 .	Non-ASCII	This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
		Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
6	Variable Length	Sequence(s) contain n's or Xaa's which represented more than one residue.
		As per the rules, each n or Xaa can only represent a single residue.
	•	Please present the maximum number of each residue having variable length and
		indicate in the (ix) feature section that some may be missing.
7	Patentin ver. 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
		sequence(s) Normally, Patentin would automatically generate this section from the
		previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section
		to the subsequent amino acid sequence. This applies primarily to the mandatory <220>-<223>
		sections for Artificial or Unknown sequences.
		Sections for Attitudes of Summer of Section 19
8	Skipped Sequences	Sequence(s) missing. If intentional, please use the following format for each skipped sequence:
·	(OLD RULES)	(2) INFORMATION FOR SEQ ID NO:X:
	(525.1522)	(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
		(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
		This sequence is intentionally skipped
		· · · · · · · · · · · · · · · · · · ·
		Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
9	Skipped Sequences	Sequence(s) missing. If intentional, please use the following format for each skipped sequence.
	(NEW RULES)	<210> sequence id number
	,	<400> sequence id number
	•	000
10	Use of n's or Xaa's	Use of n's and/or Xaa's have been detected in the Sequence Listing.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
	,	In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
11	Use of <213>Organism	Sequence(s) are missing this mandatory field or its response.
	(NEW RULES)	
12	Use of <220>Feature	Sequence(s) are missing the <220>Feature and associated headings.
	(NEW RULES)	Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
	`\\\	Please explain source of genetic material in <220> to <223> section.
	<del></del>	(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
		Toog , organizational significant and the state of the st
13	Datastia var 20 "ha"	Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted
13	Patentin ver. 2.0 "bug"	Title, Tesalting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).
	<del>-</del>	me, reading in missing mandatory numeric identifies and respondes for indicates or remoderate names.

AKS-Biotechnology Systems Branch- 5/15/99

Instead, please use "File Manager" or any other means to copy file to floppy disk.

1648 Diskette Net RAW SEQUENCE LISTING DATE: 12/26/2000 PATENT APPLICATION: US/09/497,997 TIME: 14:24:34 Corrected Diskette Needed Input. Set : A:\06600166.txt Output Set: N:\CRF3\12262000\1497997.raw 3 <110> APPLICANT: TERNYNCK, THERESA AVRAMEAS, ALEXANDRE BUTTIN, GERARD AVRAMEAS, STRAITIS SARON, MARIE-FRANCOISE BLONDEL, BRUNO 8 COUDERC, THERESA 10 MICHELSON, SUSAN 11 ZIPETO, DONATO 13 <120> TITLE OF INVENTION: VECTORS DERIVED FROM ANTIBODIES FOR TRANSFERRING SUBSTANCES INTO CELLS 15 <130> FILE REFERENCE: 0660-0166-0XCONT 17 <140> CURRENT APPLICATION NUMBER: 09/497,997 18 <141> CURRENT FILING DATE: 2000-02-04 20 <150> PRIOR APPLICATION NUMBER: PCT/FR98/01740 21 <151> PRTOR FILING DATE: 1998-08-04 Ker 1.823 d) nen Seguera Rules, the only valid (2137 responses are: Unknown, 23 <160> NUMBER OF SEO ID NOS: 35 25 <170> SOFTWARE: PatentIn version 3.0 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 17 29 <212> TYPE: PRT 30 <213> ORGANISM Artificial/Unknown Artificial Sequence, or scientific name 32 <220> FEATURE: 33 <221> NAME/KEY: misc\_feature 34 <222> LOCATION: ()..() (benus/species) 35 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide 38 <400> SEQUENCE: 1 40 Thr Arg Gln Lys Tyr Asn Lys Arg Ala Met Asp Tyr Trp Gly Gln Gly 41 1 43 Thr 46 <210> SEQ ID NO: 2 47 <211> LENGTH: 17 48 <212> TYPE: PRT 49 <213> ORGANISM Artificial/Unknown 51 <220> FEATURE: 52 <221> NAME/KEY: misc\_feature 53 <222> LOCATION: ()..() 54 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide 57 <400> SEQUENCE: 2 give source of genetic material-see circled portion of them 12 on Ever Summary Sheet 59 Thr Arg Gln Lys Tyr Gly Lys Arg Gly Met Asp Tyr Trp Gly Gln Gly 60 1 62 Thr 65 <210> SEQ ID NO: 3 66 <21.1> LENGTH: 18 67 <212> TYPE: PRT\_ 68 <213> ORGANISM: Artificial/Unknown 70 <220> FEATURE: 71 <221> NAME/KEY: misc\_feature

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PATENT APPLICATION: US/09/497,997
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                Output Set: N:\CRF3\12262000\I497997.raw
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79 1
81 Gly Thr
84 <210> SEQ ID NO: 4
85 <211> LENGTH: 22
86 <212> TYPE: PRT
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90 <221> NAME/KEY: misc_feature
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98 1
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                                        10
100 Val Lys Gly Arg Phe Thr
1.01
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1.05 <212> TYPE: PRT
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114 <400> SEQUENCE: 5
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117 1
                   5
119 Ile Lys Gly Arg Phe
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127 <220> FEATURE
128 <221> NAME/KEY: misc_feature
129 <222> LOCATION: ()..()
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133 <400> SEQUENCE: 6
135 Val. Ala Ala Ile Ser Arg Gly Gly Gly Tyr Ser Tyr Tyr Leu Asp Ser
136 1
                    5
                                         10
138 Val Lys Gly Arg Phe Thr Tle 139 \phantom{\bigg|}20\phantom{\bigg|}
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RAW SEQUENCE LISTING

DATE: 12/26/2000

TIME: 14:24:34

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/497,997

DATE: 12/26/2000

TIME: 14:24:34

RAW SEQUENCE LISTING DATE: 12/26/2000 PATENT APPLICATION: US/09/497,997 TIME: 14:24:34

Input Set : A:\06600166.txt

Output Set: N:\CRF3\12262000\1497997.raw

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/497,997

DATE: 12/25/2000

TIME: 14:24:34

DATE: 12/26/2000 TIME: 14:24:35

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/497,997

Input Set : A:\06600166.txt
Output Set: N:\CRF3\12262000\1497997.raw